

Copyright (c) 1993 - 2000 Compugen Ltd.

GenCore version 4.5

Database : SPTREMBL-17:
Maximum Match 100%
Listing first 45 summaries

```
1: sp_arachnid:*
2: sp_bacteria:*
3: sp_fungi:*
4: sp_human:*
5: sp_invertebrate:*
6: sp_mammal:*
7: sp_mhc:*
8: sp_organelle:*
9: sp_phage:*
10: sp_plant:*
11: sp_reptile:*
12: sp_virus:*
13: sp_vertebrate:*
14: sp_unclassified
```

score greater than or equal to the score of the result being printed and is derived by analysis of the total score distribution.

SUMMARIES

9dflwo	yaba-like d
9emr3	amsacta mco
97377	shigella fl
9kb44	oryza sativ
7298	plasmodium
89337	african swi
20279	rickettsia
w478	drosophila
81052	arabidopsis
65171	clostridium
3052	schizosacch
9yvvo	melanopus
CBR2	lactococcus
AZR2	bacteriopha
7517	cyanorhabdi
9WTE2	oryzias lat.
0365	caenorhabdi
95wh5	arabidopsis
Q2336	cyanorhabdi

ALIGNMENTS

Copyright (c) 1993 - 2000 Compugen Ltd.
GenCore version 4.5

20	48.5	32.1	293	2	050727	borrelia	bub
21	48.5	32.1	329	12	040852	040852	influenza
22	48.5	32.1	779	12	092f36	092f36	bradyrhizoid
23	48.5	32.1	1192	5	09W475	09W475	drosophila
24	48	31.8	275	2	051250	051250	borrelia
25	48	31.8	453	2	031020	031020	streptococci
26	47.5	31.5	148	5	09N339	09N339	caenorhabditis
27	47.5	31.5	248	5	023373	023373	caenorhabditis
28	47.5	31.5	307	5	023531	023531	caenorhabditis
29	47.5	31.5	314	5	Q18860	Q18860	caenorhabditis
30	47.5	31.5	314	5	Q9U3R4	Q9U3R4	caenorhabditis
31	47.5	31.5	314	5	Q9Y50	Q9Y50	caenorhabditis
32	47.5	31.5	329	12	040847	040847	influenza
33	47.5	31.5	339	5	021759	021759	caenorhabditis
34	47.5	31.5	339	5	001855	001855	caenorhabditis
35	47.5	31.5	339	5	017225	017225	caenorhabditis
36	47.5	31.5	339	5	017521	017521	caenorhabditis
37	47.5	31.5	339	5	044640	044640	caenorhabditis
38	47.5	31.5	339	5	Q18300	Q18300	caenorhabditis
39	47.5	31.5	339	5	Q18360	Q18360	caenorhabditis
40	47.5	31.5	339	5	017746	017746	caenorhabditis
41	47.5	31.5	339	5	Q20886	Q20886	caenorhabditis
42	47.5	31.5	339	5	Q21908	Q21908	caenorhabditis
43	47.5	31.5	339	5	Q45860	Q45860	caenorhabditis
44	47.5	31.5	339	5	Q9U359	Q9U359	caenorhabditis
45	47.5	31.5	339	5	Q9U1V7	Q9U1V7	caenorhabditis

STIC-ILL

From: Winkler, Ulrike
Sent: Tuesday, March 05, 2002 7:04 PM
To: STIC-ILL

STIC: I would really like to get a copy of the following thesis for evaluation of PCT/US01/32136.

Thanks, Ulrike

Ulrike Winkler, Ph.D.
Patent Examiner
Art Unit 1648, Mail Box 8E12
Crystal Mall 1, 8D09
1911 South Clark Place
Arlington, VA 22202
tel. 703-308-8294
fax. 703-308-4426

REFERENCE 1 (bases 1 to 144575)

AUTHORS Lee,H.J.

JOURNAL Thesis (2000) Sir William Dunn School of Pathology, University of Oxford, Oxford, UNITED KINGDOM

REFERENCE 2 (bases 1 to 144575)

AUTHORS Lee,H.J., Essani,K. and Smith,G.L.

TITLE The genome sequence of Yaba-like disease virus, a yatapoxvirus

JOURNAL Virology 281 (2), 170-192 (2001)

MEDLINE 21176366

REFERENCE 3 (bases 1 to 144575)

AUTHORS Lee,H.J.

TITLE Direct Submission

JOURNAL Submitted (13-JUL-2000) Lee H.J., Sir William Dunn School of Pathology, University of Oxford, South Parks road, Oxford, OX1 3RE, UNITED KINGDOM